Sample size calculations for comparing rate of decline

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ADC Data Core Meeting, Bethesda, MD

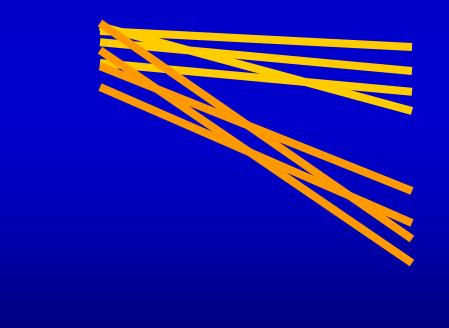
October 6, 2007

E.g., use ADC data to power rate of decline analysis for:

cohort studies
clinical trials
grants to analyze existing data (NACC proposals)

E.g., AD treatment trial Outcome: MMSE







E.g., Cohort Study (Wilson, Bennett et al. Neuroepidemiology 2006;26:61-67)

Table 2. Relation of odor identification score to baseline level of function and annual rate of change in different cognitive domains

| Cognitive domain | Model term | Estimate (SE) | p value |
|----------------------|---------------------|----------------|---------|
| Perceptual speed | Time | -0.071 (0.016) | < 0.001 |
| | Odor identification | 0.089 (0.016) | < 0.001 |
| | $Odor \times time$ | 0.015 (0.006) | 0.013 |
| Episodic memory | Time | -0.045 (0.016) | 0.004 |
| | Odor identification | 0.085 (0.012) | < 0.001 |
| | $Odor \times time$ | 0.012 (0.006) | 0.030 |
| Semantic memory | Time | -0.056 (0.014) | < 0.001 |
| | Odor identification | 0.081 (0.011) | < 0.001 |
| | $Odor \times time$ | 0.007 (0.005) | 0.156 |
| Working memory | Time | -0.049 (0.019) | 0.009 |
| | Odor identification | 0.074 (0.014) | < 0.001 |
| | $Odor \times time$ | 0.012 (0.007) | 0.084 |
| Visuospatial ability | Time | -0.007 (0.022) | 0.751 |
| | Odor identification | 0.059 (0.014) | < 0.001 |
| | $Odor \times time$ | -0.003 (0.008) | 0.667 |

Estimates are from mixed-effects models adjusted for age, sex, and education and indicate the effect of a 1-point change in odor identification score.

Possible Analytic Methods

- Least Squares 'Summary Measure'
- Random Effects Model / reml
- Marginal Model / gee

Least Squares 'Summary Measure'

- aka the NIH method
- aka 'two-stage' analysis
- Cook and Ware: "we recommend this two-stage analysis both for its efficiency and ease of interpretation." (*Annual Review of Public Health.* 1983; 4:1-23)

Least Squares 'Summary Measure'

- Esp. good for prevalent case data less prone to spurious findings (Milliken & Edland, SIM 2000) useful for describing relationship between stage of disease and rate of decline (e.g. Morris, Edland et al. Neurol 1995) Power using t-test formula
- (Schlesselman, 1971)

Power Formulas

- Random Effects Model / reml
- Marginal Model / gee

Power formula - RE model *(Hartley and Rao Biometrics;1966)

N/Arm = $2[X'V^{-1}X]_{2,2}^{-1}(z_{1-\alpha/2} + z_{1-\beta})^2 / \Delta^2$

where

- $X = (1, t) = the \ design \ matrix$
- V = Var(Y)
- \[\Lambda = detectable effect size = detectable difference in mean rate of decline
 balanced data, Var(Y) assumed known

*(Liu and Liang Biometrics;1997)

N/Arm ~ $2[XV^{-1}X]_{2,2}^{-1}(z_{1-\alpha/2} + z_{1-\beta})^2 / \Delta^2$

where

- $X = (1, t) = the \ design \ matrix$
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*Choices described for Var(Y):

unstructured
 of form σ²R, R = Cor(Y)
 compound symmetry

autoregressive

*Liu and Liang Biometrics (1997); see also Rochon SIM (1998), Jung and Ahn SIM (2003), and others

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$Var(Y) = \sigma^2 R$ implies parallel line trajectories

AD trajectories fan apart

Simulation study:

Power using AD pilot data and compound sym. assumption (ADAS-cog, Δ =1.2, power = 80% and 90%)

Simulate true power (given slopes fan apart)

 Nominal power
 (Sample Size)
 Observed Power

 80%
 (m=104)
 24%

 90%
 (m=139)
 30%

Therefore,

Use *V* = *Var*(*Y*) implied by model with random intercepts *and* random slopes:

N/Arm = $2[X'V^{-1}X]_{2,2}^{-1}(z_{1-\alpha/2} + z_{1-\beta})^2 / \Delta^2$

$V = V(Y_i) = Var(\alpha_i + \beta_i t_{ij} + \varepsilon_{ij}) = \dots$

 $V^{-1} = ...$

 $[X V^{-1}X]^{-1} = \dots$

 $N/Arm = \dots$

where

 $\sigma^2 = \sigma_{\beta}^2 + \sigma_{\epsilon}^2 / \Sigma (t - t.)^2$

where

 $\sigma^{2} = \sigma_{\beta}^{2} + \sigma_{\epsilon}^{2} / \Sigma (t - t.)^{2}$ Variance of random slopes

where

 $\sigma^2 = \sigma_{\beta}^2 + \sigma_{s}^2 / \Sigma (t - t.)^2$

Residual error variance

where

 $\sigma^2 = \sigma_\beta^2 + \sigma_s^2 / \Sigma (t - t.)^2$

Estimable by random effects model fit to pilot data

sample pilot data model fit

```
>lme(y~time, random = ~time|id)
```

```
Linear mixed-effects model fit by REML
```

```
Random effects:
Formula: ~time | id
```

```
StdDev Corr
(Intercept) 5.575794 (Intr)
time 2.382019 0.158
Residual 3.028220
```

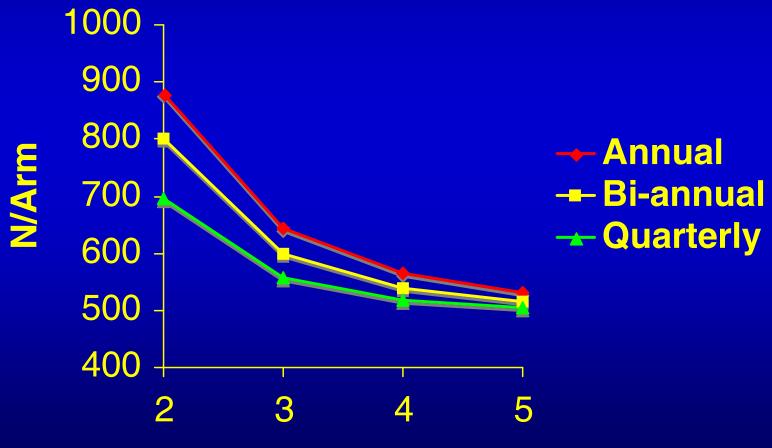
Fixed effects: y ~ time Value Std.Error DF t-value (Intercept) 16.706180 0.5945337 599 28.099634 time 1.637609 0.2642732 599 6.196652

where

 $\sigma^2 = \sigma_{\beta}^2 + \sigma_{\epsilon}^2 / \Sigma (t - t.)^2$

Determined by Study Design

N/Arm as a function of design (Alzheimer's treatment trial, outcome = ADAS-cog, effect size = 33% reduction in mean slope)



Years followup

where

 $\sigma^2 = \frac{\sigma_{\beta}^2}{\rho^2} + \frac{\sigma_{s}^2}{\Sigma(t - t.)^2}$

Varies by Instrument

Sample Size, *Prevention Trial* with Biannual Sampling, 2 or 3 Year Followup, 6 Month Sampling Interval, Effect Size = 50% Reduction in Mean Slope, Power = 90%

| | Mean | | ~ | N/Arm | |
|---------------------------|-------|------------------|----------------|-------|-----|
| | Slope | σ_{β} | σ _ε | 2Yr | 3Yr |
| Word List Delayed Rec. | 17 | 0.20 | 1.27 | 1985 | 784 |
| WMSR LM I | .73 | 1.18 | 2.44 | 595 | 354 |
| WMSR LM II | .89 | 1.20 | 2.48 | 415 | 247 |

(Pilot data courtesy OHSU ADC, Jeffery Kaye Director)



- increase N to account for expected dropout rate
- Pilot data should be representative of study population (else, see Liu and Liang 1977 for covariate weighted power formula)

Conclusions: 1

- Sample size can be dramatically underestimated when the compound symmetric model is used
- E.g., Alzheimer treatment trial setting:
 - Nominal power = 90%
 - Actual power = 30%

Conclusions: 2

The covariance structure implied by a random intercepts, random slopes model:

- is more consistent with typical longitudinal data
- can be expressed in terms of σ_{β}^2 and σ_{ϵ}^2 (easily estimated from pilot data)
- leads to heuristically appealing power formula